

SEQUENCE LISTING

SEQ ID NO:1

Length of Sequence: 19

Type of Sequence: nucleic acid

5 Number of Strand: single

Topology: linear

Type of Sequence: other type, synthetic DNA

Feature of Sequence:

KEY indicating Feature: unsure

10 Location: 1.. 19

Method of Determining Feature: E

Sequence

ACACAATCCA TTGCGAACC 19

15 SEQ ID NO:2

Length of Sequence: 20

Type of Sequence: nucleic acid

Number of Strand: single

Topology: linear

20 Type of Sequence: other type, synthetic DNA

Feature of Sequence:

KEY indicating Feature: unsure

Location: 1.. 20

Method of Determining Feature: E

25 Sequence

AGATAGTTCC TGTCTCCGCC 20

SEQ ID NO:3

Length of Sequence: 21

Type of Sequence: nucleic acid

Number of Strand: single

5 Topology: linear

Type of Sequence: other type, synthetic DNA

Feature of Sequence:

KEY indicating Feature: unsure

Location: 1.. 21

10 Method of Determining Feature: E

Sequence

CAGCTGGCAT AGCAACTAAG G 21

SEQ ID NO:4

15 Length of Sequence: 20

Type of Sequence: nucleic acid

Number of Strand: single

Topology: linear

Type of Sequence: other type, synthetic DNA

20 Feature of Sequence:

KEY indicating Feature: unsure

Location: 1.. 20

Method of Determining Feature: E

Sequence

25 GTGGTTAGCA CTGACAAGCG 20

SEQ ID NO:5

Length of Sequence: 20

Type of Sequence: nucleic acid

Number of Strand: single

Topology: linear

Type of Sequence: other type, synthetic DNA

5 Feature of Sequence:

KEY indicating Feature: unsure

Location: 1.. 20

Method of Determining Feature: E

Sequence

10 TGTTCA GTGG AACAGCAACC 20

SEQ ID NO:6

Length of Sequence: 22

Type of Sequence: nucleic acid

15 Number of Strand: single

Topology: linear

Type of Sequence: other type, synthetic DNA

Feature of Sequence:

KEY indicating Feature: unsure

20 Location: 1.. 22

Method of Determining Feature: E

Sequence

AGATTGTTCA CTGACGTCCA CC 22

25 SEQ ID NO:7

Length of Sequence: 20

Type of Sequence: nucleic acid

Number of Strand: single

Topology: linear

Type of Sequence: other type, synthetic DNA

Feature of Sequence:

KEY indicating Feature: unsure

5 Location: 1.. 20

Method of Determining Feature: E

Sequence

CTGATGCACT GCCTATGAGC 20

10 SEQ ID NO:8

Length of Sequence: 20

Type of Sequence: nucleic acid

Number of Strand: single

Topology: linear

15 Type of Sequence: other type, synthetic DNA

Feature of Sequence:

KEY indicating Feature: unsure

Location: 1.. 20

Method of Determining Feature: E

20 Sequence

CATGAGGCCT GTTGTAGAGC 20